

Mark J Abraham

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/652244/publications.pdf>

Version: 2024-02-01

9
papers

15,781
citations

1162889

8
h-index

1474057

9
g-index

10
all docs

10
docs citations

10
times ranked

20623
citing authors

#	ARTICLE	IF	CITATIONS
1	Heterogeneous parallelization and acceleration of molecular dynamics simulations in GROMACS. <i>Journal of Chemical Physics</i> , 2020, 153, 134110.	1.2	275
2	Sharing Data from Molecular Simulations. <i>Journal of Chemical Information and Modeling</i> , 2019, 59, 4093-4099.	2.5	26
3	Trends in Data Locality Abstractions for HPC Systems. <i>IEEE Transactions on Parallel and Distributed Systems</i> , 2017, 28, 3007-3020.	4.0	61
4	Tackling Exascale Software Challenges in Molecular Dynamics Simulations with GROMACS. <i>Lecture Notes in Computer Science</i> , 2015, , 3-27.	1.0	581
5	GROMACS: High performance molecular simulations through multi-level parallelism from laptops to supercomputers. <i>SoftwareX</i> , 2015, 1-2, 19-25.	1.2	14,414
6	Direct-Space Corrections Enable Fast and Accurate Lorentz-Berthelot Combination Rule Lennard-Jones Lattice Summation. <i>Journal of Chemical Theory and Computation</i> , 2015, 11, 5737-5746.	2.3	112
7	Performance enhancements for GROMACS nonbonded interactions on BlueGene. <i>Journal of Computational Chemistry</i> , 2011, 32, 2041-2046.	1.5	8
8	Optimization of parameters for molecular dynamics simulation using smooth particle-mesh Ewald in GROMACS 4.5. <i>Journal of Computational Chemistry</i> , 2011, 32, 2031-2040.	1.5	217
9	Ensuring Mixing Efficiency of Replica-Exchange Molecular Dynamics Simulations. <i>Journal of Chemical Theory and Computation</i> , 2008, 4, 1119-1128.	2.3	71